

5'-CCATCCTCTTGAAAATCTC-3' ① (SEQ ID NO:5)

5'-TCTCRTCTCACAGTTGGC-3' ② (SEQ ID NO:6).

*b7c* (6) The sixth invention is an oligonucleotide characterized in that the gene sequence of a spacer region between a gene coding 16S rRNA and a gene coding 23S rRNA of *Pectinatus cerevisiiphilus* has at least one of the following sequence group or the corresponding complementary sequence:

5'-CACTCTTACAAGTATCTAC-3' ③ (SEQ ID NO:7)

5'-CCACAATATTCCGACCAGC-3' ④ (SEQ ID NO:8)

5'-AGTCTTCTACTGCCATGC-3' ⑤ (SEQ ID NO:9).--

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Please replace the two paragraphs beginning at page 4, line 15, and ending at page 4, line 22, with the following rewritten paragraphs:

--(11) The eleventh invention is a method as in (9), wherein the nucleotide sequence coding the 16S rRNA gene of *Pectinatus frisingensis* has the following sequence:

5'-CGTATCCAGAGATGGATATT-3' ⑥ (SEQ ID NO:10)

*b7c* (12) The twelfth invention is a method as in (10), wherein the nucleotide sequence coding the 16S rRNA gene of *Pectinatus cerevisiiphilus* has the following sequence:

5'-CGTATGCAGAGATGCATATT-3' ⑦ (SEQ ID NO:11)--

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IN THE CLAIMS

Please amend claims 5, 6, 11 and 12 as follows: